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FIG. 1 - 1

Set I.

V _H 4-39			D6-13			J _H 5			
C	A	R	H/Q	C	A	R	S	D	
TGT	GCG	AGA	CA	TGT	GCG	AGA	CA	TGT	immuncytoma
C	A	S	R	G	G	G	G	AAC	Y09249
TGT	GCG	AGC	TCC	AGA	GCG	AGC	TCC	AGA	
C	A	R	H	L	G	G	G	TTC	
TGT	GCG	AGA	CAT	CTG	GCG	AGA	CAT	TTC	CLL039
C	A	R	R	F	G	G	G	TTC	CLL057
TGT	GCG	AGA	CGG	TTC	GCG	AGA	CGG	TTC	
C	A	R	S	T	G	G	G	TTC	CLL114
TGT	GCG	AGG	TCG	ACC	GCG	AGG	TCG	TTC	
C	A	R	Q	A	G	G	G	TTC	CLL202
TGT	GCG	AGA	CAA	GCT	GCG	AGA	CAA	TTC	
C	A	R	H	E	G	G	G	TTC	CLL209
TGT	GCG	AGA	CAT	GAG	GCG	AGA	CAT	TTC	

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FIG. 1 - 2

Set II.

V _H 4-34		D5-5		J _H 6	
C	A R G	V	D T A M V	Y Y Y Y	
TGT	GCG AGA GG	GTG GAT ACA	GCT ATG GTT AC	AT TAC TAC TAC	
C	A R G	Y G D T	P T I R	R Y Y Y	
TGT	GCG AGA GGA	TAC TAC	ACC ACC	ATT AGA AGA	TAC TAT CLL183
C	A R G	Y A D T	P V F R	R R Y Y	
TGT	GCG AGA GGA	TAT TAT	GCT GCT	TTT CGG CGC	TAC TAC CLL240
C	A R G	W G D T	P M L K	R R Y Y	
TGT	GCG AGA GGC	TGG TGG	ATG ATG	CTT AAA AGA	TAC TAC CLL342
C	A R A	Y P D T	P M V R	R R Y Y	
TGT	GCG AGA GCA	TAC TAC	ATG ATG	GTC AGG AGG	TAC TAC AF087422 CLL 4B 1
C	A R G	F P D T	D V I K	R R Y Y	
TGT	GCG AGA GGC	TTC TTC	GAT GAT	ATT AAG CGC	TAC TAC AJ239333 CLL ID 47 2

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FIG. 1 - 3

Set via.

V _R ¹ -02	D6-19		J _H ⁴		
C A R D/E			Y F D Y		
TGT GCG AGA GA			AC TAC TTT GAC TAC		
V . TAG	Q W L V				
GG GTA TAG CAG TGG CTG GTG C					
C A R E Q	L V L	E H Y F D Y			
TGT GCG AGC GAG CAG TGG CTG GTG	GTA	CAC TAC TTT GAC TAC		CLL011	
C A R V Q	L G L	- H F D Y			
TGT GCG AGA GTG CAG TGG CTG	GTC	--- CAC TTT GAC TAC		CLL270	
C A R E Q	L G A	- N F D Y			
TGT GCG AGA GAG CAG TGG CTG	GCA	--- AAC TTT GAC TAC		CLL266	
C A R E Q	L V L	- N F D Y			
TGT GCG AGG GAG CAG TGG CTG	GTA	--- AAC TTT GAC TAC		CLL340	
C A R V Q	L L L	- R F D Y			
TGT GCG AGA GTT CAG TGG TTA	CTC	CGA TTT GAC TAC		AJ239371	CLL 3 ²
C A R N Q	L G L	D Y F D Y			
TGT GCG AGA AAC CAG TGG CTC	GGT	GAC TAC TTT GAC TAC		AJ487492	SMZL Tierens,A.M. U3
C A R E Q	L V R	- S F D Y			
TGT GCG AGA GAG CAG TGG CTG	GTA	--- AGC TTT GAC TAC		U86787	CLL-H2B ¹ U0

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FIG. 1 - 6

Set V.

V _H 1-69		D3-10		J _H 5	
C	A R D/E			Y Y Y	
TGT	GCG AGA GA			AT TAC TAC TAC	
	I T M				
	GT ATT ACT ATG				
		V V R G V I I			
		GTT GTT CGG GGA GTT ATT ATA AC			
C	A E G M	V Q G V I		Y Y Y	
TGT	GCG GAG GGT ATG	GTT GGT GGA GTT ATT		TAC TAC TAC TAC	
C	A R S M	V Q G V I		L Y Y	
TGT	GCG AGG TCT ATG	GTT GGT GGA GTT ATT		CTC TAC TAC TAC	
C	A R A M	V R G V I		D Y Y	
TGT	GCG AGG GCT ATG	GTT GGT GGA GTT ATT		GAC TAC TAC TAC	
C	A R V M	V R G V I		L D Y Y	
TGT	GCG AGA GTT ATG	GTT GGT GGA GTT ATT		CTG GAC TAC TAC	
				AF376959	SIN Digheiro U0
				AJ389179	GN12 ⁸
				AAC51697	FUH ⁵
				AJ239372	ID64 ²

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FIG. 1 - 7

Set III.

V_R3-21
 C A R E/D
 TGT GCG AGA GA
 Y Y Y Y Y G M D V W G
 AT TAC TAC TAC TAC TAC GGT ATG GAC GTC TGG GGC

J_R6
 C A R D A N G M D V W G
 TGT GCG AGA GAT GCG AAT GGA ATG GAC GTC TGG GGC
 C A R D R N G M D V W G
 TGT GCG AGA GAT GCG AAC GGT ATG GAC GTC TGG GGC
 C A R D Q N G M D V W G
 TGT GCG AGA GAT CAA AAC GGT ATG GAC GTC TGG GGC
 C A S D R N G M D V W G
 TGT GCG AGC GAT CGA AAC GGT ATG GAC GTC TGG GGC
 C A R E P Y G M D V W G
 TGT GCG AGA GAG CCA TAC GGT ATG GAC GTC TGG GGC
 C A R D G S G M D V W G
 TGT GCG AGA GAT GGC TCC GGT ATG GAC GTC TGG GGC
 C A R D A N G M D V W G
 TGT GCG AGA GAT GCT AAC GGC ATG GAC GTC TGG GGC

CLL282
 CLL175
 CLL412
 ID28²
 AJ239379
 AF174100 sc77u-16⁹
 AJ389178 GN11⁸
 AF299104 IF case 10¹⁰ U3

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FIG. 1 - 8

Set Vie		J _H 4		Y F D Y		AC TAC TTT GAC TAC	
V _H 5-51 D6-19		C A R Q/H		TGT GCG AGA CA		GG GTA TAG CAG TGG CTG GTA C	
C A R Q		C A R Q		C A R Q		C A R Q	
TGT GCG AGG	CAG CAG	Q Q	W W	L L	G G	D D	Y Y
C A R R	Q Q	W W	L L	G G	GGT GGT	TAC TAC	TTT TTT
TGT GCG AGA	AGG CAG	Q Q	W W	L L	A A	G G	H H
C A R R	Q Q	W W	L L	G G	CTA GGC	CAC TTT	GAC TAC
TGT GCG AGA	CAG CAA	Q Q	W W	L L	G G	Y Y	F F
C A R R	Q Q	W W	L L	G G	TTC GGC	TAC TAC	TTT TTT
TGT GCG AGA	CAG CAG	Q Q	W W	L L	V V	P P	Y Y
C A R R	Q Q	W W	L L	G G	CTT CTT	TAC TAC	TTT TTT
TGT GCG AGA	GAG CAG	Q Q	W W	L L	I I	T T	H H
C A R R	Q Q	W W	L L	G G	ATA ATA	ACT ACT	CAC TTT
TGT GCG AGA	CAG CAG	Q Q	W W	L L	V V	L L	D D
C A R R	Q Q	W W	L L	G G	CTG CTG	TAC TAC	TTT TTT
TGT GCG AGA	CAG CAG	Q Q	W W	L L	V V	L L	S S
C A R R	Q Q	W W	L L	G G	Not available		

PH1562 HOW¹³

Tre¹¹ O12/0-2 JK2

CLL026

AF099198

AJ414007

AJ239373

AJ555263

AJ272398

AG¹²

PH1562

HOW¹³

CLL021 Russia U0

ID38²

G014¹

FIG. 1 - 9

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V _H 1-02										D6-19					J _H 4				
C	A	R	D/E												F	D	Y		
TGT	GCG	AGA	GA												AC	TTT	GAC	TAC	
				V	Q	W	L	V											
	GG	GTA	TAG	CAG	TGG	CTG	GTA	C											
C	A	R	E	Q	W	L	V	L	E	H	Y	F	D	Y					
TGT	GCG	AGG	GAG	CAG	TGG	CTG	GTA	CTT	GAG	CAC	TAC	TTT	GAC	TAC	CLL011				
C	A	R	V	Q	W	L	G	L	R	-	H	F	D	Y					
TGT	GCG	AGA	GTG	CAG	TGG	CTG	GCC	TTA	AGA	---	CAC	TTT	GAC	TAC	CLL270				
C	A	R	E	Q	W	L	G	A	E	-	N	F	D	Y					
TGT	GCG	AGA	GAG	CAG	TGG	CTG	GGC	GCA	GAA	---	AAC	TTT	GAC	TAC	CLL266				
C	A	R	E	Q	W	L	V	L	K	-	N	F	D	Y					
TGT	GCG	AGG	GAG	CAG	TGG	CTG	GTA	CTG	AAA	---	AAC	TTT	GAC	TAC	CLL340				
C	A	R	V	Q	W	L	L	L	E	-	R	F	D	Y					
TGT	GCG	AGA	GTT	CAG	TGG	TTA	TTA	CTC	GAA	---	CGA	TTT	GAC	TAC	AJ239371	CLL 3 ²			
C	A	R	N	Q	W	L	G	L	D	-	Y	F	D	Y					
TGT	GCG	AGA	AAC	CAG	TGG	CTG	GGT	CTC	GAC	---	TAC	TTT	GAC	TAC	AJ487492	SMZL Tierens, A.M.			
C	A	R	E	Q	W	L	V	R	T	-	S	F	D	Y					
TGT	GCG	AGA	GAG	CAG	TGG	CTG	GTA	AGG	ACG	---	AGC	TTT	GAC	TAC	U86787	CLL-H2B ¹ U0			
C	A	R	E	Q	W	L	V	L	S	-	Y	F	D	Y					
TGT	GCG	AGG	GAG	CAG	TGG	CTG	GTC	CTA	TCT	---	TAC	TTT	GAC	TAC	CLL336				
C	A	R	E	Q	W	L	V	L	-	N	Y	F	D	Y					
TGT	GCG	AGG	GAG	CAG	TGG	CTG	GTA	CTT	---	AAC	TAC	TTT	GAC	TAC	CLL360				
C	A	R	E	Q	W	L	A	L	K	-	P	F	D	Y					
TGT	GCG	AGA	GAG	CAG	TGG	CTG	GCC	TTA	AAA	---	CCC	TTT	GAC	TAC	AF376961	LAN Digheiro			
C	A	R	K	Q	W	L	A	I	V	N	Y	F	D	Y					
TGT	GCG	AGA	AAG	CAG	TGG	CTG	GCC	ATC	GTC	AAC	TAC	TTT	GAC	TAC	L01278	CLL-412 ³			
C	A	R	E	Q	W	L	G	L	P	-	T	F	D	Y					
TGT	GCG	AGA	GAG	CAG	TGG	CTG	GGT	CTA	CCT	---	ACC	TTT	GAC	TAC	U86801 ⁴	U3/1			
C	A	R	V	Q	W	L	G	L	T	G	P	N	D	Y					
TGT	GCT	AGG	GTT	CAG	TGG	CTG	GGC	CTG	ACG	GGG	CCG	AAT	GAC	TAC	U84176	KEM (VH1-46) ⁵ U0			
C	A	R	G	Q	W	L	V	I	L	-	N	F	D	Y					
TGT	GCG	AGG	GGA	CAG	TGG	CTG	GTC	ATC	CTA	---	AAC	TTT	GAC	TAC	U84162	BYR (VH1-46) ⁵ U0			
C	A	R	D	Q	W	L	P	T	-	N	N	F	D	Y					
TGT	GCG	AGA	GAT	CAG	TGG	CTG	CCC	ACG	---	AAC	AAC	TTT	GAC	TAC	AF376953	Digheiro PIQ U2			
C	A	R	E	Q	W	L	V	L	S	-	H	F	D	Y					
TGT	GCG	AGG	GAG	CAG	TGG	TTG	GTA	CTA	TCT	---	CAC	TTT	GAC	TAC	CLL154	(VH1-18)			
C	A	R	Q	Q	W	L	G	G		D	Y	F	D	Y					
TGT	GCG	AGG	CAG	CAG	TGG	CTG	GGT	GGC		GAC	TAC	TTT	GAC	TAC	CLL026				
C	A	R	R	Q	W	L	A	L		G	H	F	D	Y					
TGT	GCG	AGA	AGG	CAG	TGG	CTG	GCC	CTA		GGC	CAC	TTT	GAC	TAC	AF099198	Tre ¹¹ O12/0-2			
JK2																			
C	A	R	Q	Q	W	F	G	V		Y	Y	F	D	Y					
TGT	GCG	AGA	CAG	CAA	TGG	TTC	GGC	GTG		TAC	TAC	TTT	GAC	TAC	AJ414007	CLL021 Russia U0			
C	A	R	Q	Q	W	L	V	L		P	Y	F	D	Y					
TGT	GCG	AGA	CAG	CAG	TGG	CTG	GTA	CTT		CCA	TAC	TTT	GAC	TAC	AJ239373	ID38 ²			
C	A	R	E	Q	W	L	I	V		T	H	F	D	Y					
TGT	GCG	AGA	GAG	CAG	TGG	CTC	ATA	GTA		ACT	CAC	TTT	GAC	TAC	AJ555263	G014			
C	A	R	Q	Q	W	L	V	L		D	Y	F	D	Y					
TGT	GCG	AGA	CAG	CAG	TGG	CTG	GTG	TTG		GAC	TAC	TTT	GAC	TAC	AJ272398	AG ¹²			
C	A	R	E	Q	W	L	V	L		S	N	F	D	Y					
Not available										PH1562					HOW ¹³				

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FIG. 2

Amino acid alignments of the H chain variable regions of all sequences in each Set

Set	IV	CDR1	CDR2	CDR3
Germline	QVQLVQSGAEVKKFGSSVKVSKASGGTSSSYAISWVRQAPGQGLEWNGGIIPFGTANYAQKFGQGRVTITADKSTSTAYNELSSLRSEDTAVYYCARD	YDYVWGSYRY	DAFDVWGQGTMTVSS	
MCLL068
MCCL258
MF9
SHI
CLL022
RIZ
Consensus	QVQLVQSGAEVKKFGSSVKVSKASGGTSSSYAISWVRQAPGQGLEWNGGIIPFGTANYAQKFGQGRVTITADKSTSTAYNELSSLRSEDTAVYYCARD	YDYVWGSYRY	DAFDVWGQGTMTVSS	
Set	VIII	CDR1	CDR2	CDR3
Germline	QVQLVQSGAEVKKFGSSVKVSKASGGTSSSYAISWVRQAPGQGLEWNGGIIPFGTANYAQKFGQGRVTITADKSTSTAYNELSSLRSEDTAVYYCARD	DIVVPAAI	YIFYGNDVWGQGTMTVSS	
FS41
UCM4
G013
Consensus	QVQLVQSGAEVKKFGSSVKVSKASGGTSSSYAISWVRQAPGQGLEWNGGIIPFGTANYAQKFGQGRVTITADKSTSTAYNELSSLRSEDTAVYYCARD	DIVVPAAI	YIFYGNDVWGQGTMTVSS	
Set	V	CDR1	CDR2	CDR3
Germline	QVQLVQSGAEVKKFGSSVKVSKASGGTSSSYAISWVRQAPGQGLEWNGGIIPFGTANYAQKFGQGRVTITADKSTSTAYNELSSLRSEDTAVYYCARD	DMVRGVI	YIFYGNDVWGQGTMTVSS	
CLL561
M321
RF22
GRI2
FUH
ID64
SIN
Consensus	QVQLVQSGAEVKKFGSSVKVSKASGGTSSSYAISWVRQAPGQGLEWNGGIIPFGTANYAQKFGQGRVTITADKSTSTAYNELSSLRSEDTAVYYCARD	DMVRGVI	YIFYGNDVWGQGTMTVSS	
Set	II	CDR1	CDR2	CDR3
Germline	QVQLQQWAGLLKPKSETLSLTCAVYGESFSGYTHSWIRQPPGKGLWNGIINPNSGGTNYAQKFGQGRVTITADKSTSTAYNELSSLRSEDTAVYYCARD	QVDTAMV	YIFYGNDVWGQGTMTVSS	
CLL183
CLL240
CLL342
CLL48
ID47
Consensus	QVQLQQWAGLLKPKSETLSLTCAVYGESFSGYTHSWIRQPPGKGLWNGIINPNSGGTNYAQKFGQGRVTITADKSTSTAYNELSSLRSEDTAVYYCARD	QVDTAMV	YIFYGNDVWGQGTMTVSS	
Set	Via	CDR1	CDR2	CDR3
Germline	QVQLVQSGAEVKKFGASVKVSKASGYTFTSYAMHWVRQAPGQGLEWNGIINPNSGGTNYAQKFGQGRVTITADKSTSTAYNELSSLRSEDTAVYYCARD	QVDTAMV	YIFYGNDVWGQGTMTVSS	
CLL011
CLL266
CLL270
CLL340
CLL3
CLL18
CLL-H2B
Consensus	QVQLVQSGAEVKKFGASVKVSKASGYTFTSYAMHWVRQAPGQGLEWNGIINPNSGGTNYAQKFGQGRVTITADKSTSTAYNELSSLRSEDTAVYYCARD	QVDTAMV	YIFYGNDVWGQGTMTVSS	
Set	Vib	CDR1	CDR2	CDR3
Germline	QVQLVQSGAEVKKFGASVKVSKASGYTFTSYAMHWVRQAPGQGLEWNGIINPNSGGTNYAQKFGQGRVTITADKSTSTAYNELSSLRSEDTAVYYCARD	QVDTAMV	YIFYGNDVWGQGTMTVSS	
CLL336
CLL360
DAN
CLL-412
POR
PIQ
Consensus	QVQLVQSGAEVKKFGASVKVSKASGYTFTSYAMHWVRQAPGQGLEWNGIINPNSGGTNYAQKFGQGRVTITADKSTSTAYNELSSLRSEDTAVYYCARD	QVDTAMV	YIFYGNDVWGQGTMTVSS	
Set	Vic	CDR1	CDR2	CDR3
Germline	QVQLVQSGAEVKKFGASVKVSKASGYTFTSYAMHWVRQAPGQGLEWNGIINPNSGGTNYAQKFGQGRVTITADKSTSTAYNELSSLRSEDTAVYYCARD	QVDTAMV	YIFYGNDVWGQGTMTVSS	
CLL154
Set	Vid	CDR1	CDR2	CDR3
Germline	QVQLVQSGAEVKKFGASVKVSKASGYTFTSYAMHWVRQAPGQGLEWNGIINPNSGGTNYAQKFGQGRVTITADKSTSTAYNELSSLRSEDTAVYYCARD	QVDTAMV	YIFYGNDVWGQGTMTVSS	
BYR
KSH
Consensus	QVQLVQSGAEVKKFGASVKVSKASGYTFTSYAMHWVRQAPGQGLEWNGIINPNSGGTNYAQKFGQGRVTITADKSTSTAYNELSSLRSEDTAVYYCARD	QVDTAMV	YIFYGNDVWGQGTMTVSS	
Set	Vie	CDR1	CDR2	CDR3
Germline	EVQLVQSGAEVKKFGESLRISCKGSGISFTSYMISWVRQMPGKGLWNGRIDPSDSTNYSPFQGHVTSADKSIISTAYLQNSSLKASDTAMYYCARD	QVDTAMV	YIFYGNDVWGQGTMTVSS	
CLL026
TR8
CLL021
ID36
G014
AG
HOW
Consensus	EVQLVQSGAEVKKFGESLRISCKGSGISFTSYMISWVRQMPGKGLWNGRIDPSDSTNYSPFQGHVTSADKSIISTAYLQNSSLKASDTAMYYCARD	QVDTAMV	YIFYGNDVWGQGTMTVSS	

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FIG. 3

Amino acid alignments of the L chain variable regions of all sequences in each Set.

set IV
 Germline EIVLTQSPGTLSPGERATLSCRASQSVSSYLAWYQQKPGAPRLIYQASRATGIPDRFSGSGGTFTLTISRLEPEDFATYCCQYGSFPTFGQGTKEIK CDR3
 CLL068
 CLL1258
 MF9
 Consensus EIVLTQSPGTLSPGERATLSCRASQSVSSYLAWYQQKPGAPRLIYQASRATGIPDRFSGSGGTFTLTISRLEPEDFATYCCQYGSFPTFGQGTKEIK L.....G.....

set VIII
 Germline EIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGAPRLIYDASNRATGIPARFSGSGGTFTLTISRLEPEDFATYCCQSRNTFGGTKEVDIK CDR3
 G013
 Consensus EIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGAPRLIYDASNRATGIPARFSGSGGTFTLTISRLEPEDFATYCCQSRNTFGGTKEVDIK G.....

set V -
 Germline QSVLTQPPASGTPGQRTVITSCGSSNIGNTVWYQQLEGTAPKLLIYNNQRPFGVPDRFSGSKSGTSASIAISGLQSEDEADYYCAWDDSLNGPVFGGKLTVL CDR2 CDR3
 RF22
 GN12
 Consensus QSVLTQPPASGTPGQRTVITSCGSSNIGNTVWYQQLEGTAPKLLIYNNQRPFGVPDRFSGSKSGTSASIAISGLQSEDEADYYCAWDDSLNGPVFGGKLTVL R.....F.....

set I
 Germline DVVMTQSPFLSLPVTIGQFASISCRSSQSLVDSGNTYLNWYQQRPQSPRLIYKVSRDSGVDRFSGSGGTFTLTISRVEADVGYYICMGTHTPPYTFQGTKEIK CDR2 CDR3
 CLL183
 CLL240
 CLL342
 Consensus DVVMTQSPFLSLPVTIGQFASISCRSSQSLVDSGNTYLNWYQQRPQSPRLIYKVSRDSGVDRFSGSGGTFTLTISRVEADVGYYICMGTHTPPYTFQGTKEIK L.....L.....L.....M.....V.....

set VI
 Consensus DIQMTQSPSLASVGDRTVITTCRASQSISSYLWYQQKPGAPKLLIYAASISQSGVSRFSGSGGTFTLTISLQPEDFATYCCQSYSTPP WTFGQGTKEIK CDR1 CDR2 CDR3
 CLL011
 CLL266
 CLL270
 CLL340
 CLL336
 CLL360
 CLL-412
 CLL154
 TRF
 G014
 Consensus DIQMTQSPSLASVGDRTVITTCRASQSISSYLWYQQKPGAPKLLIYAASISQSGVSRFSGSGGTFTLTISLQPEDFATYCCQSYSTPP -YTFGQGTKEIK Y.....L.....L.....R.....L.....

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FIG. 5

D2-2																JH6	VRL6	Jk3
CARD/E																Y Y Y	Q Q R S N W P P	
tgt gcg aga ga																at tac tac tac	Cag cag cgt agc aac tgg cct cc	
D I V V P A I																		F T F G P G
ag gat att gta gta cca gct gct ata cc																		a ttc act ttc ggc cct ggg
G G M S - . .																		
. gg ggc g t.c																		
G A M G - . .																		
G G M R - . .																		
. gc gg g .ga																		
1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18																		
1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18																		

V _H 1-69	D3-10	J _H 6	J _λ 1/3*
C A R D/E	Y Y Y	V _λ 1-16	W D D S L N G P
tgt gcg aga ga	at tac tac tac		tgg gat gac agc ctg aat ggt cc
I T M V R G V I I			Y V F G
gt att act atg gtt cgg gga gtt att ata ac			t tgg gtg ttc ggc
A Q Q T CLL561			
A Q T Y MJ21			
G Q T Y RF22			
E G Q G I GN12			
S Q N V L FUH			
A Q H L D ID64			
V Q S L D SIN			

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FIG. 7

V _H 4-34	D5-5	J _H 6	V _K A17	J _K 1/2*
C A R G	Y Y Y	M Q G T H W P P		
tgt gcg aga gg	at tac tac tac	atg caa ggt aca cac tgg cct cc		
V D T A M V				W T F G
gtg gat aca gct atg gtt ac				g tgg acg ttc ggc
Y G . . . P T I R R . . . CLL183				
. a tac .g. c. cc a. ga aga t				
Y A . . . P V F R R . . . CLL240				
. a tat .c. t c. g. t. egg cg.				
W G . . . P . L K R . . . CLL342				
. c tgg .g. c. c. a aga				
A Y P . . . P . V R R . . . CLL 4B				
. ca tac cc. c. c.c .gg agg				
F P . . . D V I K R . . . ID47				
. c ttc cc. a. g. a. g cg.				
1 2 3 4 5 6 7 8 9 10 11 12 13 14 15			1 2 3 4 5 6 7 8 9 10 11 12	

V _H 1-02										D6-19										J _H 4										V _K 012/02										J _K 1/2*									
CARD/E										Y F D Y										Q Q S Y S T P P																													
tgt gcg aga ga										ac tac ttt gac tac										caa cag agt tac agt acc cct cc																													
V * Q W L V																														W T F G																			
gg gta tag cag tgg atg gta a																														g tgg acg tta gga																			
. L E H CLL011									 T																													
. G A E - N CLL266									 - K																													
. V G L R - H CLL270									 -																													
. L K - N CLL340									 -																													
. V L L E - R CLL3									 -																													
. N G L D slv18 (marginal zone lymphoma)									 -																													
. E R T - S CLL-H2B									 -																													
. L S - CLL336									 S -																													
. L - N CLL360									 -																													
. A L K - P LAN									 -																													
. K A I V N CLL-412									 -																													
. P T - N N PIQ									 -																													
. L S - H CLL154									 -																													
. G I L - N BYR									 -																													
. V G L T G P N KEM									 -																													
. G G - D CLL026									 -																													
. R A L - G H TRE									 -																													
. F G V - Y CLL021									 -																													
. L P - ID38									 -																													
. E I V T - H GO14									 - R																													
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